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AMENDMENT: IN THE SPECIFICATION

Please enter the Sequence Listing into the application.

1. Please replace the paragraph [0018] with the following amended paragraph(s):

1A. Agonist-induced downregulation of opioid receptors. Cells stably expressing MOR, DOR or D MOR (~1-2 pmol/mg) were treated with agonist (10 μM DADLE, an opioid peptide having the sequence Tyr-D-Ala-Gly-Phe-D-Leu; SEQ ID NO:7; Sigma, St. Louis, MO) for 3 hours or left untreated. Cells were then chilled on ice, washed extensively and total opioid radioligand binding sites were determined for each cell line (16). Both DOR and D MOR showed significant downregulation whereas MOR was not substantially downregulated under the same conditions (p<.001). Error bars represent s.d. from a representative experiment (n=3 experiments), with each data point derived from triplicate determinations.

2. Please replace the paragraph [0113] with the following amended paragraph(s):

[0113] The invention also provides GASP polypeptides. A GASP polypeptide of the invention includes a GASP amino acid sequence, i.e., an amino acid sequence that has at least about 70% identity to GASP SEQ ID NO:2 (GASP1) or GASP SEQ ID NO:6 (GASP2) over a comparison window of at least 15 contiguous amino acids. GASP SEQ ID NO:2 (GASP1) is the amino acid sequence of a GASP polypeptide described in detail in Example 1. The nucleic acid and (single-letter code) amino acid sequences of these polypeptides are given below.

GASP1 Nucleic Acid Sequence

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ctatattgcatctagttctggttctgaggatgagtctgttaagacaccctggttctgggccagagataaaaccaatacctggtctgggcccagg gaag at ccca at ag cagg tcc agg ttt agg tct aagaa ag aag tct at gtt gaat ca ag ttct gg at ct gag cat gaag accatt tg gag tcct agg at cagg tcc agg tct agg at cagg tcc agg tct agg at cagg tcc agg tccggtttggggctggaaaggaggcaaattcaggtccaaaatgagagctgggaaggaggccaataacagggccaggcaaggccaagc gagaagcttg cattgattt catgcctgggt ctatagatgtaattaaaaaagagtcctgtttctggcctgaagaaaatgctaatacctttt caaggccagg t cag agg agg agg context t gg gacct gg tt ctg gg ctacag ac gag t cag cat gg cag at gaag ccag cat agag t cag tt ctg gat cag ac gag tag agg cag at gaag ccag cat agag t cag tag agg cag at gag agg cag at gag at gaga caagtgg agg at gag te cata at tgg gag tt gg tt ctg gae tgaag aa gag ge cag tat gg gg ac tgg gg ctag cag taa at ceagac cag tat gag gag tag gag tagcattggtccctggttttggtctggagaacaagttgatatagaggctggaatcggagaagaggccaggccaggagctgaagaagaagacaatattcgggtcctggttttgggctgaaaaccagacctatatggattgtagggctgaaactagctgtgacaccatgcaaggggctgaggaggag gagcccattattgggtcctggttttggaccagagtagaagcttgtgtggagggtgatgtcaacagcaagtctagcctggaggacaaggaag gaggacattgtcaattcgtggttctggtccagaaaatacacaaagccagaggccattatagggtcctggttatgggctacagaagagagtaa tatagatgggactggagaaaaggccaagttactgactgaagaggagaccataatcaattcctggttctggaaagaagatgaagccatttcag cattagaagagaggctgggtcttgcagcaaatccagtcctaaagctgaagaggaagaagtcattattgggtcctggttctgggaagaagag gccagtccggaggcagtggcaggagtcggctttgagtcaaagcctgggactgaggaggaagaaatcactgttgggtcctggttctggcct gaagaagaagccagtatacaggctggatctcaggcagtagaggaaatggagtcagagactgaagaggaaaccatttttgggtcctggttctgggatggaaaagaagtcagtgaagaagcaggaccatgctgtgtatccaagccagaggatgatgaagaagatgattgttgagtcctggttctggtctagagacaaagccattaaggaaactggaactgtggccacctgtgagtccaagccagaaaatgaggaaggggccattgttgggtcttg gtttgaggctgaagatgaggtagataacaggactgacaatggaagcaactgtgggtccaggacattagctgatgaagatgaggccatagtg gggtcctggttctgggcaggagatgaggcccattttgaatcaaatcctagccccgtgttcagggccatttgcaggtccacgtgttcagttgaa caggag acct gate cttcacg caggact cagagt tgggaggagg teact gttcagt teag acct ggtccat ggggt tagg ttcccat

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GASP1 Amino Acid Sequence

YWTFFVTCTMTGAEIESGAQVKPEKKPGEEVVGGAEIENDVPLVVRPKVRTQAQIMPG ARPKNKSKVMPGASTKVETSAVGGARPKSKAKAIPVSRFKEEAQMWAQPRFGAERLSK TERNSQTNIIASPLVSTDSVLVAKTKYLSEDRELVNTDTESFPRRKAHYQAGFQPSFRSK EETNMGSWCCPRPTSKQEASPNSDFKWVDKSVSSLFWSGDEVTAKFHPGNRVKDSNRS MHMANQEANTMSRSQTNQELYIASSSGSEDESVKTPWFWARDKTNTWSGPREDPNSRS RFRSKKEVYVESSSGSEHEDHLESWFGAGKEGKFRSKMRAGKEANNRARHRAKREACI DFMPGSIDVIKKESCFWPEENANTFSRPMIKKEARARAMTKEEAKTKARARAKQEARSE EEALIGTWFWATDESSMADEASIESSLQVEDESIIGSWFWTEEEASMGTGASSKSRPRTD GERIGDSLFGAREKTSMKTGAEATSESILAADDEQVIIGSWFWAGEEVNQEAEEETIFGS WFWVIDAASVESGVGVSCESRTRSEEEEVIGPWFWSGEQVDIEAGIGEEARPGAEEETIF GSWFWAENOTYMDCRAETSCDTMQGAEEEEPIIGSWFWTRVEACVEGDVNSKSSLEDK **EEAMIPCFGAKEEVSMKHGTGVRCRFMAGAEETNNKSCFWAEKEPCMYPAGGGSWKS** RPEEEEDIVNSWFWSRKYTKPEAIIGSWLWATEESNIDGTGEKAKLLTEEETIINSWFWK EDEAISEATDREESRPEAEEGDIVGSWFWAGEEDRLEPAAETREEDRLAAEKEGIVGSWF GAREETIRREAGSCSKSSPKAEEEEVIIGSWFWEEEASPEAVAGVGFESKPGTEEEEITVG SWFWPEEEASIQAGSQAVEEMESETEEETIFGSWFWDGKEVSEEAGPCCVSKPEDDEEM IVESWFWSRDKAIKETGTVATCESKPENEEGAIVGSWFEAEDEVDNRTDNGSNCGSRTL ADEDEAIVGSWFWAGDEAHFESNPSPVFRAICRSTCSVEQEPDPSRRPQSWEEVTVQFKP GPWGRVGFPSISPFRFPKEAASLFCEMFGGKPRNMVLSPEGEDQESLLQPDQPSPEFPFQ YDPSYRSVQEIREHLRAKESTEPESSSCNCIQCELKIGSEEFEELLLLMEKIRDPFIHEISKI AMGMRSASQFTRDFIRDSGVVSLIETLLNYPSSRVRTSFLENMIRMAPPYPNLNIIQTYIC KVCEETLAYSVDSPEQLSGIRMIRHLTTTTDYHTLVANYMSGFLSLLATGNAKTRFHVL KMLLNLSENLFMTKELLSAEAVSEFIGLFNREETNDNIQIVLAIFENIGNNIKKETVFSDD DFNIEPLISAFHKVEKFAKELQGKTDNQNDPEGDQEN

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GASP2 Nucleic Acid Sequence

at gat gt ccctct ggt ggt cag accca ag gt tag gaccca gg caactact gg gg caa gg ccca aa act gag acca ag tct gt gcct gc gg according to the contract graph of the contract graph ocaaggcccaaaactgaggcccaagcaatgtctggggcaaggcccaaaactgaggtccaagtaatgggtggtgcaagacccaaaacgga ggctcaaggaatcacaggggccaggcccaaaaccgatgccagggcagtaggtggcgctcgttctaaaactgatgccaaggcaatccctg gagcaaggcccaaggatgaggcccaggcatgggcccagagtgaatttgggactgaagcagtgtcacaggcagaaggagtgtcccaga ctaatgccgttgcttggccactggccactgctgagtctggatcagttactaaatctaagggcctgtctatggatagagaactagtcaatgtgga tgctgaaacctttcctggcacccagggtcagaaaggaatccagccctggtttggaccaggggaggagactaatatggggtcttggtgctatt agagacaagtgtcagatcatggcccagggaagagtccaataccaggtccaggcacagggctaaacatcagactaatcccaggtccaggc ccaga tcca agca agaa gcctat gtt gattcct ggtct ggatct gaggat gaggccag ccagca acccattct ccttct gggtt ggagaa aataccalled a comparent of the comparent gattch gattch gaggat gaggac gaggaa gccag gaggagaa acccattct gggtt ggagaa aataccalled gaggat gaggac gaggac gaggaa gagaa gaggac gaggaa gaggac gaggaa gagaa gaggac gaggaa gagaa gaggac gaggaa gagaa gagaaataacttgttcaggcccagagtcagggagggggaaatatcaggtccaagctcaggacaaatagagaagattgttttgaatctgagtctgaa gatgagttctataagcagtcctgggttttgcctggagaagaggccaatagtagattcaggcacagagacaaagaagatcctaatactgccttgcagaaaaagaggccagtttggagggtggagcttcagcaatctgtgaatctgagccaggaactgaggagggggccattggcggatccgc gtactgggctgaggaaaagtccagtttgggggctgtggccagagaagaggccaagccggagtctgaagaagaggccatatttgggtcctggttctgggacagagatgaggcctgctttgacctaaatccctgtcctgtgtacaaggtcagtgataggttcagagatgcagctgaggagctta atgcatcctccaggccccaaacctgggacgaggtcactgttgaattcaaacctggtctttttcatggggttggcttccgatccacaagcccctt tggaattcccgaagaggcttctgaaatgcttgaggcaaagcccaagaacctggaacttagcccagaaggagaagagcaggaatctttgctt cagcetgatcagcetagtcetgagttcacatttcagtatgatcettcetaceggtcagtcegggaaattcgagagcatcttagggccagggag agtgcagagtctgagagttggtcctgcagctgcatacaatgtgagctgaaaattggttctgaagagtttgaagaattccttttattaatggacaa aattegggateettttatteatgaaatatetaaaattgeaatgggtatgagaagtgetteteaatttaeeegagattteattegagatteaggtgttg totcact tattgaaaacctt gettaat tatccatect ctag ag ttag gacaa g ttttt tggaaaa tatgat teacat g gete cacet tatccaa at ctaal tattag aa acctt gettaat tatccatect ctag ag ttag gacaa g ttttt tggaaaa tatgat teacat g gete cacet tatccaa at ctaal tattag aa acctt g ctaal tattag aaacatgattgagacattcatatgtcaagtgtgtgaggaaacccttgcacatagtgtggattcccttgagcagctgactggaataaggatgcttag a cacct cact at gact at tgact at cacacac tgat tgccaact at at gtccgggtttctctcct tat taaccacagc caat gcgagaac gaag tto the catter of the cacacac transfer of the cacacacac transfer of the cacacacac transfer of the cacacacac transfer of the cacacacac transfer of the cacacac transfer of the cacacacatagaagagacaaatgataatattcaaattgttattaaaatgtttcagaatatcagtaacattataaaaagtggaaagatgtccttaattgatga tgatttcagtcttgagccgcttatttctgcatttcgtgaatttgaggagttagctaagcaactacaagcccaaatagaccacaaatgatcctga ggtgggacaacaaagttaa

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GASP2 Amino Acid Sequence

MTGAEIEPSAQAKPEKKAGEEVIAGPERENDVPLVVRPKVRTQATTGARPKTETKSVPA
ARPKTEAQAMSGARPKTEVQVMGGARPKTEAQGITGARPKTDARAVGGARSKTDAKA
IPGARPKDEAQAWAQSEFGTEAVSQAEGVSQTNAVAWPLATAESGSVTKSKGLSMDRE
LVNVDAETFPGTQGQKGIQPWFGPGEETNMGSWCYSRPRAREEASNESGFWSADETST
ASSFWTGEETSVRSWPREESNTRSRHRAKHQTNPRSRPRSKQEAYVDSWSGSEDEASNP
FSFWVGENTNNLFRPRVREEANIRSKLRTNREDCFESESEDEFYKQSWVLPGEEANSRFR
HRDKEDPNTALKLRAQKDVDSDRVKQEPRFEEEVIIGSWFWAEKEASLEGGASAICESE
PGTEEGAIGGSAYWAEEKSSLGAVAREEAKPESEEEAIFGSWFWDRDEACFDLNPCPVY
KVSDRFRDAAEELNASSRPQTWDEVTVEFKPGLFHGVGFRSTSPFGIPEEASEMLEAKPK
NLELSPEGEEQESLLQPDQPSPEFTFQYDPSYRSVREIREHLRARESAESESWSCSCIQCEL
KIGSEEFEEFLLLMDKIRDPFIHEISKIAMGMRSASQFTRDFIRDSGVVSLIETLLNYPSSRV
RTSFLENMIHMAPPYPNLNMIETFICQVCEETLAHSVDSLEQLTGIRMLRHLTMTIDYHT
LIANYMSGFLSLLTTANARTKFHVLKMLLNLSENPAVAKKLFSAKALSIFVGLFNIEETN
DNIQIVIKMFQNISNIIKSGKMSLIDDDFSLEPLISAFREFEELAKQLQAQIDNQNDPEVGQ
QS

SEQ ID NO:6

These amendments are made without prejudice and are not to be construed as abandonment of the previously claimed subject matter or agreement with any objection or rejection of record.